

Human GIP (inactive)



www.mesoscale.com®

Ordering Information

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Scientific Support

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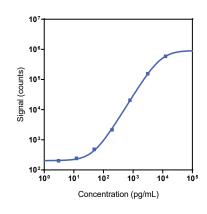
Company Address

Meso Scale Discovery A division of Meso Scale Diagnostics, LLC. 1601 Research Boulevard Rockville, MD 20850-3173 USA

Product Options	Catalog Number	Description			
Multiplex	K151ACM, K251ACM	U-PLEX Metabolic Group 1 (human)			
Singleplex	K1515TK-1/-2/-4	U-PLEX Human GIP (inactive) Assay with SECTOR™ plates			
	K1515TK-21/-22/-24	U-PLEX Human GIP (inactive) Assay with QuickPlex Ultra™ plates			
	K2515TK-2/-4	U-PLEX Human GIP (inactive) Assay with 384-well plates			
Antibody Set	B215T-2/-3	U-PLEX Human GIP (inactive) Antibody Set			
Protocol	U-PLEX Product Inserts are available at www.mesoscale.com				

The MESO SCALE DISCOVERY® U-PLEX platform was designed to provide ultimate flexibility for detection of biomarkers in a wide variety of sample types. This datasheet provides the representative performance of the U-PLEX® Human GIP (inactive) Assay tested on U-PLEX 96-well SECTOR plates run as a multiplex. The data do not represent the product specifications. Under your experimental conditions, the assay may perform differently from the representative data. U-PLEX assays are offered in either singleplex or multiplex; both are available on 96- or 384-well plates. See a U-PLEX product insert for instrument compatibility.

Representative Calibration Curve and Sensitivity



Assay	Median LLOD (pg/mL)	LLOD Range (pg/mL)		
GIP (inactive)	27	22-39		

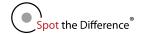
The Calibrator curve was fitted with a 4-parameter logistic model with a $1/Y^2$ weighting. The lower limit of detection (LLOD) is a calculated concentration corresponding to 2.5 standard deviations above the background (zero Calibrator).

Precision

Control	Average Conc. (pg/mL)	Average Intra-run Conc. (%CV)	Inter-run Conc. (%CV)
High	2,560	6.6	12.0
Mid	2,000	7.7	23.3
Low	1,530	5.9	16.4

Controls were made by spiking Calibrator into assay diluent at 3 levels within the quantitative range of the assay. Average intra-run concentration %CV is the average %CV of the control replicates within an individual run. Inter-run concentration %CV is the variability of controls across multiple runs.

For Research Use Only. Not for use in diagnostic procedures.





MSD® U-PLEX Human GIP (inactive)

Tested Samples

Sample Type	Serum (N=12)	EDTA Plasma (N=12)	P800 Plasma (N=8)		
Median (pg/mL)	85	154	117		
Range (pg/mL)	ND-121	ND-154	ND-212		
% Detected	25	8	50		

Normal serum, EDTA plasma, and P800 plasma samples were diluted 4-fold prior to the assay. ND = non-detectable (<LLOD)

Dilution Linearity

Serum			EDTA Plasma			P800 Plasma			Cell Culture Media		
Fold Dilution	Average % Recovery	% Recovery Range	Fold Dilution	Average % Recovery	% Recovery Range	Fold Dilution	Average % Recovery	% Recovery Range	Fold Dilution	Average % Recovery	% Recovery Range
2	67	48-80	2	79	67-93	2	77	67-93	2	92	82-99
8	112	109-116	8	104	98-111	8	108	98-117	8	99	90-110
16	116	108-122	16	102	86-115	16	112	102-126	16	104	91-123

Normal human serum, EDTA plasma, P800 plasma, and cell culture media were spiked with Calibrator and tested at different dilutions. Percent recovery at each dilution level was normalized to the dilution-adjusted, 4-fold concentration. Samples may benefit from additional dilution with assay diluent to reduce matrix effects.

% Recovery = (measured concentration / expected concentration) x 100

Spike Recovery

	Serum		EDTA Plasma		P800 Plasma		Cell Culture Media	
Spike Level	Average % % Recovery Recovery Range		Average % % Recovery Range		Average % Recovery			% Recovery Range
High	65	60-74	87	81-103	74	64-79	88	80-103
Mid	67	58-75	81	75-100	74	66-83	85	79-89
Low	71	62-82	81	71-86	76	68-84	87	82-95

Normal serum, EDTA plasma, P800 plasma, and cell culture media were spiked with Calibrator at 3 levels. Spiked samples were diluted 4-fold to determine the expected concentration of the analyte. Samples may benefit from additional dilution with assay diluent to reduce matrix effects.

% Recovery = (measured concentration / expected concentration) x 100

Specificity

To assess specificity, the GIP (inactive) Antibody Set was tested individually against a larger panel of analytes for nonspecific binding (BDNF, C-Peptide, CTACK, Desghrelin, ENA-78, Eotaxin, Eotaxin-2, Eotaxin-3, EP0, FGF-21, FGF-23, FLT3L, Fractalkine, FSH, G-CSF, Ghrelin (Ser3-octanoylated), GIP (1–42), GIP (3–42), GLP-1 (7–36), GLP-1 (9–36), Glucagon, GM-CSF, GR0- α , I-309, IFN- α 2a, IFN- β , IFN- γ , IL-1 α , IL-1 β , IL-1RA, IL-2, IL-2R α , IL-3, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IL-10, IL-12/IL-23p40, IL-12p70, IL-13, IL-15, IL-16, IL-17A/F, IL-17C, IL-17D, IL-17E/IL-25, IL-17F, IL-18, IL-21, IL-22, IL-23, IL-27, IL-29/IFN- α 1, IL-31, IL-31, IL-33, Insulin, IP-10, Leptin, LH, MCP-1, MCP-2, MCP-4, M-CSF, MDC, MIF, MIP-1 α , MIP-1 α , MIP-5, PP, Proinsulin, PYY (3–36), SDF-1 α , TNF- α , TN

% Nonspecificity = (nonspecific signal / specific signal) x 100

GIP (1-42; active) cross-reacts 2.4% with the GIP (inactive) assay. We do not recommend multiplexing the GIP (inactive) assay with the GIP (active) or GIP (total) assays on the same plate.

Diluent Compatibility

The data included in this document were collected with Assay Diluent 13 (supplemented with 1,000 klU/mL Aprotinin [provided] and 100 μ M diprotin A [not provided]) and Antibody Diluent 11. MSD offers a range of assay and antibody diluents for separate purchase. Depending on your assay needs, other diluents may be tested. Diprotin A should be purchased separately.

Assay Components

Calibrator: GIP (inactive) is included in Calibrator 13. The human GIP (inactive) Calibrator is a synthetic peptide.

Antibodies: the U-PLEX® Human GIP (inactive) Assay uses a mouse monoclonal antibody for capture and a mouse monoclonal antibody for detection.

Assay generation: A

Note: This datasheet contains representative assay performance data. In custom multiplex formats, the assay may perform differently from the representative data shown.



